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AJ245709 Homo sapi AF124141 Homo sapi AF135794 Homo sapi AX026530 Sequence

AX026529 HSA245709 AF124141

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REFERENCE AUTHORS TITLE JOURNAL SOURCE ORGANISM VERSION KEYWORDS DEFINITION ACCESSION RESULT AX026529 FEATURES Locus AX026529 Sequence AX026529 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 1547)

Masure, S.L. and Richardson, A.

Human akt-3

Patent: WO 0037613-A 1 29-JUN-2000;

MASURE STEFAN LEO JOZEF (BE); RICHARDSON ALAN (BE); JANSSEN PHARMACEUTICA NV (BE) human. Homo sapiens AX026529.1 Н Location/Qualifiers /organism="Homo sapiens" from Patent W00037613. GI:10187717 .1547 1547 bp PAT 16-SEP-2000

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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1 (bases 1 to 1547)

Masure.S., Haefner,B., Wesselink,J.J., Hoefnagel,E., Mortier,E., Verhasselt,P., Tuytelaars,A., Gordon,R. and Richardson,A.

Molecular cloning, expression and characterization of the human serine/threonine kinase Akt-3

Eur. J. Biochem. 265 (1), 353-360 (1999)
                                                                                        Masure, S.L.
Direct Submission
Submitted (25-AUG-1999)
                                                                          High-Throughput Screening, Janss
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348 g 408 t 0; Score 1547; Pred. No. 0; Mismatches DB 0; 9; Length Indels 0; Gaps 540 600 600 540 480 480 420 420 360 300 300 240 180 60 60 360 240 180 0 RESULT AF124141 LOCUS QΥ ₽ δÃ Qy Qy δÃ Qy В Q οy Qy γQ δÃ Qy 맑 Qy B δÃ 밁 Ъ Вþ Βğ Ъ B В Вb B Вb Вþ Š 1501 1501 1441 1381 1381 1261 1201 1141 1141 1021 1261 1201 1081 1081 1021 961 961 901 841 721 661 901 721 781 781 ω ttcagggctcttgataaaggatccaaataaacgccttggtggaggaccagatgatgcaaa ggactgcatggacaatgagaggcggccgcatttccctcaattttccttactctgcaagtgg gcttgtacctcctfttaaacctcaagtaacatctgagacagatactagatattttgatga gttagaagataatgactatggccgagcagtagactggtggggcctaggggttgtcatgta GGACTGCATGGACAATGAGAGGCGGCCGCATTTCCCTCAATTTTCCTACTCTGCAAGTGG AGAATTTACAGCTCAGACTATTACAATAACACCACCTGAAAAATATGATGAGGATGGTAT agaatttacagctcagactattacaataacaccacctgaaaaatatgatgaggatggtat GCTTGTACCTCCTTTTAAACCTCAAGTAACATCTGAGACAGATACTAGATATTTTGATGA GTTAGAAGATAATGACTATGGCCGAGCAGTAGACTGGTGGGGGCCTAGGGGTTGTCATGTA AATGCTGGACAAAGATGGCCACATAAAAATTACAGATTTTGGACTTTGCAAAGAAGGAT aatgctggacaaagatggccacataaaaattacagattttggacttttgcaaagaagggat AF124141 Homo sapiens AF124141 AF124141.1 G protein 1708 ďď kinase mRNA B gamma mRNA,

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Direct Submission
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99194749
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Hon
1 (bases 1 to 2811)
Nakatani,K., Sakaue,H., Thompson,D.A., Weigel,R.J. and F
Identification of a human Akt3 (protein kinase B gamma)
contains the regulatory serine phosphorylation site
Biochem. Biophys. Res. Commun. 257 (3), 906-910 (1999)
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d, Stanford, CA 94305, U
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Eukaryota; Metazoa; Chordata; C
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1 (bases 1 to 1436)
Masure, S.L. and Richardson, A.
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/db_xref="taxon:9606"
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Homo sapiens Chordata; Crani
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Mammada; Eutheria; Primates; Catar
Chases 1 to 1570)
Guo,K., Ivashchenko,Y. and Clark,K.
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AX056819
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Homo sapiens
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           Metazoa: Chordata; Craniata; Vertebrata; Euteleostomi;
Eutheria; Primates; Catarrhini; Hominidae; Homo.
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VDLPYLNNFSVAKCQLMKTERPKPNTFIIRCLOWTTVIETFIFHYDTFDEREEWTEAI
QAVADRLORDEERMNCSPTSQIDIGEEMAJSTTHHKRKTWNDEDYLKLLGKGTEG
KVILVREKASGKYYAMKILKKEVIIAKDEVAHTLTESRVLKNTRHPFLTSLKYSFQTK
DRLCFYMEYVNGGELFFHLSRERVFSEDRTRFYGAEIVSALDYLHSGKIVRDLKLEN
LMLDKDGHIKITDFGLÓKEGITDAATMKTFCGTPEYLAPEVLEDNDYGRAVDWWGLGV
VMYEMMCGRLPFYNQDHEKLFELILMEDIKFPRTLSSDAKSLLSGLLXKDDNRKLIGG
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1 (bases to 1584)
Wiemann 8. Wellenreuther, R., Gassenhuber, J., Glas
Ansorge W., Boecher, M., Bloecker, H., Bauersachs, S., Blum, H.
Lauber, J., Duesterhoeft, A., Beyer, A., Koehrer, K., Strack, N.
Mowes, H.W., Ottenwaelder, B., Obermaler, B., Tampe, J., Heubne
Mambutt, R., Korn, B., Klein, M. and Poustka, A.
Toward a Catalog of Human Genes and Proteins: Sequencing an
                                                                     Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                          Homo
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                                                                    Craniata; Vertebrata; Catarrhini; Hominidae;
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                                                                     Hominidae;
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                                                                              Euteleostomi;
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This clone (DKFZp434N0250) is available at the RZPD in This clone (DKFZp434N0250) is available at the RZPD in Please contact the RZPD: Ressourcenzentrum, Heubnerweg Berlin-Charlottenburg, GERMANY; Email: clone@rzpd.de Fu information about the clone and the sequencing project at http://www.mips.biochem.mpg.de/proj/cDNA/.
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Submitted (15-SEP-1999) MIPS, Am Klopferspitz 18a, D-82152
Martinsried, GERMANY
Clone from S. Wiemann, Molecular Genome Analysis, German Car
Research Center (DKFZ); Email s.wiemann@dkfz-heidelberg.de;
sequenced by DKFZ (German Cancer Research Center,
Heidelberg/Germany) within the cDNA sequencing consortium of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Poustka, A.,
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vdlpyplymissyakcolmkterpkpyptpilircpythudtperermyteai
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kvilymekasckyyamkilkkevilakdevahtltesrvlkwrrhpfitslkysgfotk
kvilymekasckyyamkilkrevirakdevahtltesrvlkwrrhpfitslkysgfotk
drlgtvmeyvnggelffhlsrervfsedrtrfygaeivsaldylhsgkivyrdiklen
lmjdrdghikitdfglckegftdaatwktfcctpeylapevledndygravdwwgglgv
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DH10B; sites NotI + SalI"
/dev_stage="adult"
154. .1542
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/db_xref="gL/5912043"
/db_xref="5MISS-PROT:09Y243"
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                                           agaaattatgagacacagtttcttctctggagtaaactggcaagatgtatatgataaaaa
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Two Splice Variants of Protein Kinase B gamma
Regulatory Capacity Depending on the Presence
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LMLDKNOHIKTTDFGLCKEGTTDAATMKTFCGTPEYLAPEVLEDNDYGRAVDWWGLGV
VMYEMMCGRLPFYNQDHEKLFELILMEDIKFPRTLSSDAKSLLSGLLIKDPNKRLGGG
PDDAKEIMRHSFFSGVNWQDVYDKKLVPPFKPQVTSETDTRYFDEEFTAQTITITPPE
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/gene="AKT3"
/note="PKBgammal; C-terminal splice variant lacking the Ser472 phosphorylation site; alt
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/db_xref="GI:15072340"
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/db_xref="taxon:9606"
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74. .1471
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Pred. No. 4.3e-304;
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Direct Submission
Submitted (27-JAN-1999) Friedrich Miescher-Institut,
Maulbeerstrasse 66, Basel 4058, Switzerland
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                                                                                                                                                                                                                                                                                                                    Molecular cloning and characterization of a new member of the protein kinase family: association of the pleckstrin homology domain of three types of RAC protein kinase with protein kinas subspecies and beta gamma subunits of G proteins Biochem. Biophys. Res. Commun. 216 (2), 526-534 (1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Direct Submission
Submitted (23-MAR-1995) to the DDBJ/EMBL/GenBank data
Konishi, Kobe University, Biosignal Research Center;
Rokkoudai-cho 1-1, Kobe, Hyogo 657, Japan
(E-mall:hkonishi@inherit.biosig.kobe-u.ac.jp, Tel:078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rat mRNA for D49836
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Mammalia; Eutheria; Rodentia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rattus norvegicus
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Kameyama,K., Haga,T. and Kikkawa,U
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LMLDKDGHIKITDFGLCKEGITDAATMKTFCGTPEYLAPEVLEDNDYGRAVDWWGLGV
VMYEMWCGRLPFYNQDHEKLFELILWEDIKFPRTLSSDAKSLLSGLLIKDPNKRLGGG
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OAVADRLQROEEERNMCSPTSQIDNIGEEENMASTTHKKTWNDEDYLKLLGKGTEK
KVILVREKASGKYYAMKILKKEVIIAKDEVAHTLTESRVLKNTRHPFLTSLKYSFQTK
                                                                                                                                                                                                                     /db_xref="taxon:10116"
/tissue_type="brain"
47. .1411
                                                                                                                                                         /product="RAC-PK gamma"
/protein_id="BAA08637.1"
/db_xref="GI:1401040"
                                       PDDPKEIMRHSFFSGVNWQDVYDKKLVPPFKPQVTSETDTRYFDEEFTAQTITITPPE
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                                                                                           GATGCTAGATAAGGATGGCCATATAAAAATTACGGATTTTGGACTTTGCAAAGAAGGGAT
                                                                                                  aatgctggacaaagatggccacataaaaattacagattttggactttgcaaagaagggat
                                                                                                                                                                  GTCGAGAGAGCGGGTGTTCTCTGAGGACCGCACACGTTTCTATGGTGCAGAAATTGTCTC
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Query Match
Best Local Similarity
Matches 1106; Conserv

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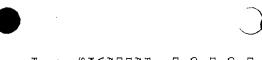
Score 850.6; DB 5; Pred. No. 1.1e-185; Mismatches



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BASE COUNT
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Phasianinae; Gallus.
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Verkleij, A.J., Hanafusa, H. and
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           PPDQDDSMDCVDNERRPHFPQFSYSASGTA"
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                                         GGGPDDAKEIMQHKFFAGIVWQDVYGKKLVPPFKPQVTSETDTRYFDEEFTAQMITII
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/protein_id="AAB94767.1"
/db_xref="GI:2745889"
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Amphibia; Batrachia; Anura;
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/product="Akt"
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/db_xref="taxon:8355"
/dev_stage="oocyte"
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                                                                                                                                                                                                                                                                                                             /note="PH-domain containing
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Batrachia; Anura; Mesobatrachia; Pipoidea;
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                                         CTAATGCTGGACAAAGACGGACACATAAAGATAACAGATTTTGGACTGTGCAAAGAAGGA
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                                                                                GCTTTAGATTATCTTCATTCTGAGAAAAATGTAGTTTACAGAGACTTGAAGTTGGAAAAT
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/translation="MGQTVTTPLSLTLEHWGDVQRIASNQSVDVKKRRWVTFCSAEWP
TFGVCMPQDGTFNLDIILQVKSKVFSBGPHGHEDQVPYIVTWEALAYEEPPBWYKEFVS
PKLSLSPTAPILPSGPSTQPPPRSALYPALTPSIKPRPSKPQVLSDNGGPLIDLLTED
PPPYGEQGPSSSDGDGDREEATSTPEIPAPSPMVSRLRGKRDPPAAVSTTSRAFPLRL
                                                             /product="gag:akt fusion
/protein_id="AAA42545.1"
/db_xref="GI:210068"
                                                                                                                                               /gene="v-akt"
324. .2615
                                                                                                                  /codon_start=1
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/db_xref="taxon:12438"
/cell_line="95785"
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LKRGEESTMDFRAGGSSONSGAEEMEVSLAK PKHRVTWNEFEYLKLLGAGTGTGKVILV
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KDGHIKITDEGLCKEGIKAGATWATGCGTPEYLLAPEVLEDNDVGRAVDWMGLGVWYK
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M.musculus |
X65687
                                                                     Submitted (24 APR.1992) A. Bellacosa, Fox Chase C Burholme Ave, Philadelphia PA 19111, USA 2 (bases 1 to 2626)
Bellacosa, A., Franke, T.F., Gonzalez-Portal, M.E., Taguchi, T., Gardner, J., Cheng, J.O., Testa, J.R. ar Structure, expression and chromosomal mapping of relationship to v-akt and its implications Oncogene 8 (3), 745-754 (1993)
                                                                Oncogene
93173519
                                                                                                                                                                                                                                                           protein kinase; serine/threonine house mouse.
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                                                                                                                                                                                 Direct Submission
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/organism="Mus musculus"
/strain="AKR/j"
/db_xref="taxon:10090"
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Mus musculus protein
M94335
M94335.1 GI:200641
M94335.1 kinase.
Mus musculus CDNA to
Mus musculus
                         and C kinases) protein kinase Unpublished
                                                              Eukaryota; Metazoa; Chordata; Mammalia; Eutheria; Rodentia; 1 (bases 1 to 1513)
Bousquets, X. and Powell, C.T.
                                                     Complete nucleotide coding
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musculus protein
Location/Qualifiers
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THDRLCFVMEYANGGELFFHLSRERVFSEDRAFEYGAEIVSALDYLSEKNVYRDLK
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GGGSEDAKEIMOHFKLFELILMEEIAFPTKPDEFTAQMITIT
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Best Local Similarity 71...
Matches 1041; Conservative
           48.3%;
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Score 746.6; DB Pred. No. 1e-161; 0; Mismatches 39
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aagaggggagaata              AAACGAGGGGAATA	8 atcatgagcgatgttaccattgtgaaagaaggttgggttcagaagaggggaggaatatata 67	20

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601	542 attctgaagaaagaagtcattattgcaaaggatgaagtggcacacactctaactgaaagc 601	Оу 54
559	500 ACCTTTGGGAAGGTGATTCTGGTGAAAGAGAAGGCCACACGCCGCTACTATGCCATGAAG 559	Db 50
541	acttttgggaaagttattttggttcgagagaaggcaagtggaaaatactatgctatgaag	Qy 482
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481	425 catcataaaagaaagacaatgaatttttgactatttgaaactactaggtaaaggc 481	Оу 42
439	380 TTCCGATCAGGCTCACCCAGTGACAACTCAGGGGCTGAAGAGATGGAGGTGTCCCTGGCC 439	Db 38
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364	305 acagaagctatccaggctgtagcagacagactgcagagggcaagaagaggaggagagaatgaat	Оу 30

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